OIPE

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/924,125

DATE: 12/13/2001
TIME: 13:58:26

Input Set : A:\ES.txt

Output Set: N:\CRF3\12132001\I924125.raw

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3 <110> APPLICANT: Communi, Didier
      5 <120> TITLE OF INVENTION: THE NATURAL LIGAND FOR ORPHAN G PROTEIN COUPLED RECEPTOR
GPR86 AND
             METHODS OF USE
      6
     8 <130> FILE REFERENCE: 9049/2092
    10 <140> CURRENT APPLICATION NUMBER: 09/924,125
    11 <141> CURRENT FILING DATE: 2001-07-08
    13 <150> PRIOR APPLICATION NUMBER: US 09/924,125
    14 <151> PRIOR FILING DATE: 2001-07-08
    16 <160> NUMBER OF SEQ ID NOS: 9
                                                                     ENTERED
     18 <170> SOFTWARE: PatentIn version 3.1
     20 <210> SEQ ID NO: 1
     21 <211> LENGTH: 1002
     22 <212> TYPE: DNA
     23 <213> ORGANISM: Homo sapiens
     25 <400> SEQUENCE: 1
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     28 atagtacage tggtattece agecetetae acagtggttt tettgacegg cateetgetg
                                                                              120
     30 aatactttgg ctctgtgggt gtttgttcac atccccagct cctccacctt catcatctac
                                                                              180
                                                                              240
     32 ctcaaaaaca ctttggtggc cgacttgata atgacactca tgcttccttt caaaatcctc
     34 tetgaeteae acetggeaee etggeagete agagettttg tgtgtegttt ttetteggtg
                                                                              300
     36 atattttatg agaccatgta tgtgggcatc gtgctgttag ggctcatagc ctttgacaga
                                                                              360
     38 ttcctcaaga tcatcagacc tttgagaaat atttttctaa aaaaacctgt ttttgcaaaa
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     40 acggtctcaa tcttcatctg gttctttttg ttcttcatct ccctgccaaa tatgatcttg
                                                                              480
     42 agcaacaagg aagcaacacc atcgtctgtg aaaaagtgtg cttccttaaa ggggcctctg
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     44 gggctgaaat ggcatcaaat ggtaaataac atatgccagt ttattttctg gactgttttt
                                                                              600
     46 atcctaatgc ttgtgtttta tgtggttatt gcaaaaaaag tatatgattc ttatagaaag
     48 tccaaaagta aggacagaaa aaacaacaaa aagctggaag gcaaagtatt tgttgtcgtg
                                                                              720
     50 gctgtcttct ttgtgtgttt tgctccattt cattttgcca gagttccata tactcacagt
                                                                              780
     52 caaaccaaca ataagactga ctgtagactg caaaatcaac tgtttattgc taaagaaaca
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     54 actototttt tggcagcaac taacatttgt atggatccct taatatacat attottatgt
                                                                              900
     56 aaaaaattca cagaaaagct accatgtatg caagggagaa agaccacagc atcaagccaa
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     58 gaaaatcata gcagtcagac agacaacata accttaggct ga
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    63 <212> TYPE: PRT
    64 <213> ORGANISM: Homo sapiens
    66 <400> SEQUENCE: 2
    68 Met Asn Thr Thr Val Met Gln Gly Phe Asn Arg Ser Glu Arg Cys Pro
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                                            10
     69 1
    72 Arg Asp Thr Arg Ile Val Gln Leu Val Phe Pro Ala Leu Tyr Thr Val
                                                            30
                    20
     73
    76 Val Phe Leu Thr Gly Ile Leu Leu Asn Thr Leu Ala Leu Trp Val Phe
                                                        45
                                    40
     77
                35
    80 Val His Ile Pro Ser Ser Ser Thr Phe Ile Ile Tyr Leu Lys Asn Thr
                                55
    81
    84 Leu Val Ala Asp Leu Ile Met Thr Leu Met Leu Pro Phe Lys Ile Leu
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75

-80

70

85 65



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88 Ser Asp Ser His Leu Ala Pro Trp Gln Leu Arg Ala Phe Val Cys Arg
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                                             90
     89
     92 Phe Ser Ser Val Ile Phe Tyr Glu Thr Met Tyr Val Gly Ile Val Leu
                                         105
                    100
     96 Leu Gly Leu Ile Ala Phe Asp Arg Phe Leu Lys Ile Ile Arg Pro Leu
                                                          125
                115
                                     120
     97
     100 Arg Asn Ile Phe Leu Lys Lys Pro Val Phe Ala Lys Thr Val Ser Ile
                                                       140
     101
             130
                                  135
     104 Phe Ile Trp Phe Phe Leu Phe Phe Ile Ser Leu Pro Asn Met Ile Leu
                                                   155
     105 145
                              150
     108 Ser Asn Lys Glu Ala Thr Pro Ser Ser Val Lys Lys Cys Ala Ser Leu
                                              170
     1.09
                          165
     112 Lys Gly Pro Leu Gly Leu Lys Trp His Gln Met Val Asn Asn Ile Cys
                                                               190
     113
                      180
                                          185
     116 Gln Phe Ile Phe Trp Thr Val Phe Ile Leu Met Leu Val Phe Tyr Val
                                      200
                                                           205
                 195
     117
     120 Val Ile Ala Lys Lys Val Tyr Asp Ser Tyr Arg Lys Ser Lys Ser Lys
                                  215
                                                       220
     121
             210
     124 Asp Arg Lys Asn Asn Lys Lys Leu Glu Gly Lys Val Phe Val Val Val
                                                   235
                              230
     128 Ala Val Phe Phe Val Cys Phe Ala Pro Phe His Phe Ala Arg Val Pro
     129
                                              250
                                                                   255
                          245
     132 Tyr Thr His Ser Gln Thr Asn Asn Lys Thr Asp Cys Arg Leu Gln Asn
     133
                                          265
                                                               270
                      260
     136 Gln Leu Phe Ile Ala Lys Glu Thr Thr Leu Phe Leu Ala Ala Thr Asn
                                                           285
     137
                 275
                                      280
     140 Ile Cys Met Asp Pro Leu Ile Tyr Ile Phe Leu Cys Lys Lys Phe Thr
                                  295
                                                       300
     141
             290
     144 Glu Lys Leu Pro Cys Met Gln Gly Arg Lys Thr Thr Ala Ser Ser Gln
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     145 305
     148 Glu Asn His Ser Ser Gln Thr Asp Asn Ile Thr Leu Gly
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     153 <211> LENGTH: 11
     154 <212> TYPE: DNA
C--> 155 <213> ORGANISM: Artificial
     157 <220> FEATURE:
     158 <221> NAME/KEY: misc_binding
     159 <222> LOCATION: (1)..(11)
     160 <223> OTHER INFORMATION: NF-kB binding element
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     164 ggggactttc c
     167 <210> SEQ ID NO: 4
     168 <211> LENGTH: 31
     169 <212> TYPE: DNA
C--> 170 <213> ORGANISM: Artificial
     172 <220> FEATURE:
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11

W--> 173 <221> NAME/KEY: primer

174 <222> LOCATION: (1)..(31)

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175 <223> OTHER INFORMATION: GPR86 human receptor: a sense primer 178 <400> SEQUENCE: 4 179 ccggaattca ccatgaacac cacagtgatg c 31 182 <210> SEQ ID NO: 5 183 <211> LENGTH: 31 184 <212> TYPE: DNA C--> 185 <213> ORGANISM: Artificial 187 <220> FEATURE: W--> 188 <221> NAME/KEY: primer 189 <222> LOCATION: (1)..(31) 190 <223> OTHER INFORMATION: GPR86 human receptor: anti-sense primer 193 <400> SEQUENCE: 5 194 cttgtctaga tcagcctaag gttatgttgt c 31 197 <210> SEO ID NO: 6 198 <211> LENGTH: 20 199 <212> TYPE: DNA C--> 200 <213> ORGANISM: Artificial 202 <220> FEATURE: W--> 203 <221> NAME/KEY: primer 204 <222> LOCATION: (1)..(20) 205 <223> OTHER INFORMATION: GPR86 sense primer 208 <400> SEQUENCE: 6 209 tgtgtcgttt ttcttcggtg 20 212 <210> SEQ ID NO: 7 213 <211> LENGTH: 18 214 <212> TYPE: DNA C--> 215 <213> ORGANISM: Artificial 217 <220> FEATURE: W--> 218 <221> NAME/KEY: primer 219 <222> LOCATION: (1)..(18) 220 <223> OTHER INFORMATION: GPR86 antisense primer 223 <400> SEQUENCE: 7 224 ctgccaaaaa gagagttg 18 227 <210> SEQ ID NO: 8 228 <211> LENGTH: 20 229 <212> TYPE: DNA C--> 230 <213> ORGANISM: Artificial 232 <220> FEATURE: W--> 233 <221> NAME/KEY: primer 234 <222> LOCATION: (1)..(20) 235 <223> OTHER INFORMATION: aldolase sense primer 238 <400> SEQUENCE: 8 239 ggcaagggca tcctggctgc 20 242 <210> SEQ ID NO: 9 243 <211> LENGTH: 23 244 <212> TYPE: DNA C--> 245 <213> ORGANISM: Artificial

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W--> 248 <221> NAME/KEY: primer

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249 <222> LOCATION: (1)..(23)

250 <223> OTHER INFORMATION: aldolase antisense reverse primer

253 <400> SEQUENCE: 9

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23

VERIFICATION SUMMARY

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Input Set : A:\ES.txt

Output Set: N:\CRF3\12132001\I924125.raw

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